

Serial Number: 001658, 699**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically:
-
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____.
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically:
-
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:
-
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:
-
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included:
-
- ☐ Deleted extra, invalid, headings used by an applicant, specifically:
-
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____.
- ☒ Inserted mandatory headings, specifically: <220> feature (Segs. #2 and 4)
- ☐ Corrected an obvious error in the response, specifically:
-
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically:
-
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____
-
-
-

*Examiner: ~~The~~ The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95

OIPE

RAW SEQUENCE LISTING DATE: 09/27/2000
 PATENT APPLICATION: US/09/658,699 TIME: 09:59:25

Input Set : A:\Cpg.pto
 Output Set: N:\CRF3\09272000\I658699.raw

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3 <110> APPLICANT: De Waal Malefyt, Rene
4       Kastelein, Robert A.
5       Lira, Sergio A.
6       Narula, Satwant
7       Oppmann, Birgit
8       Rennick, Donna M.
9       Wiekowski, Maria
11 <120> TITLE OF INVENTION: Mammalian Cytokines; Related Reagents and Methods
13 <130> FILE REFERENCE: DX01042X US
C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/658,699
C--> 16 <141> CURRENT FILING DATE: 2000-09-08
18 <150> PRIOR APPLICATION NUMBER: 09/393,090
19 <151> PRIOR FILING DATE: 1999-09-09
21 <150> PRIOR APPLICATION NUMBER: 60/164,616
22 <151> PRIOR FILING DATE: 1999-11-10
24 <160> NUMBER OF SEQ ID NOS: 5
26 <170> SOFTWARE: PatentIn Ver. 2.1
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29 <211> LENGTH: 570
30 <212> TYPE: DNA
31 <213> ORGANISM: Unknown Organism
33 <220> FEATURE:
34 <223> OTHER INFORMATION: Description of Unknown Organism: surmised Homo
35     sapiens
37 <220> FEATURE:
38 <221> NAME/KEY: CDS
39 <222> LOCATION: (1)..(567)
41 <220> FEATURE:
42 <221> NAME/KEY: mat_peptide
43 <222> LOCATION: (64)..(567)
45 <400> SEQUENCE: 1
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48   -20               -15               -10
50 gct cag ggc aga gct gtg cct ggg ggc agc agc cct gcc tgg act cag      96
51 Ala Gln Gly Arg Ala Val Pro Gly Gly Ser Ser Pro Ala Trp Thr Gln
52   -5               -1    1               5               10
54 tgc cag cag ctt tca cag aag ctc tgc aca ctg gcc tgg agt gca cat      144
55 Cys Gln Gln Leu Ser Gln Lys Leu Cys Thr Leu Ala Trp Ser Ala His
56   15               20               25
58 cca cta gtg gga cac atg gat cta aga gaa gag gga gat gaa gag act      192
59 Pro Leu Val Gly His Met Asp Leu Arg Glu Glu Gly Asp Glu Glu Thr
60   30               35               40
62 aca aat gat gtt ccc cat atc cag tgt gga gat ggc tgt gac ccc caa      240
63 Thr Asn Asp Val Pro His Ile Gln Cys Gly Asp Gly Cys Asp Pro Gln
64   45               50               55
66 gga ctc agg gac aac agt cag ttc tgc ttg caa agg atc cac cag ggt      288

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RAW SEQUENCE LISTING DATE: 09/27/2000
 PATENT APPLICATION: US/09/658,699 TIME: 09:59:26

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70 ctg att ttt tat gag aag ctg cta gga tcg gat att ttc aca ggg gag 336
71 Leu Ile Phe Tyr Glu Lys Leu Leu Gly Ser Asp Ile Phe Thr Gly Glu
72 80 85 90
74 cct tct ctg ctc cct gat agc cct gtg gcg cag ctt cat gcc tcc cta 384
75 Pro Ser Leu Leu Pro Asp Ser Pro Val Ala Gln Leu His Ala Ser Leu
76 95 100 105
78 ctg ggc ctc agc caa ctc ctg cag cct gag ggt cac cac tgg gag act 432
79 Leu Gly Leu Ser Gln Leu Leu Gln Pro Glu Gly His His Trp Glu Thr
80 110 115 120
82 cag cag att cca agc ctc agt ccc agc cag cca tgg cag cgt ctc ctt 480
83 Gln Gln Ile Pro Ser Leu Ser Pro Ser Gln Pro Trp Gln Arg Leu Leu
84 125 130 135
86 ctc cgc ttc aaa atc ctt cgc agc ctc cag gcc ttt gtg gct gta gcc 528
87 Leu Arg Phe Lys Ile Leu Arg Ser Leu Gln Ala Phe Val Ala Val Ala
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108 Ala Gln Gly Arg Ala Val Pro Gly Gly Ser Ser Pro Ala Trp Thr Gln
109 -5 -1 1 5 10
111 Cys Gln Gln Leu Ser Gln Lys Leu Cys Thr Leu Ala Trp Ser Ala His
112 15 20 25
114 Pro Leu Val Gly His Met Asp Leu Arg Glu Glu Gly Asp Glu Glu Thr
115 30 35 40
117 Thr Asn Asp Val Pro His Ile Gln Cys Gly Asp Gly Cys Asp Pro Gln
118 45 50 55
120 Gly Leu Arg Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile His Gln Gly
121 60 65 70 75
123 Leu Ile Phe Tyr Glu Lys Leu Leu Gly Ser Asp Ile Phe Thr Gly Glu
124 80 85 90
126 Pro Ser Leu Leu Pro Asp Ser Pro Val Ala Gln Leu His Ala Ser Leu
127 95 100 105
129 Leu Gly Leu Ser Gln Leu Leu Gln Pro Glu Gly His His Trp Glu Thr
130 110 115 120
132 Gln Gln Ile Pro Ser Leu Ser Pro Ser Gln Pro Trp Gln Arg Leu Leu
133 125 130 135
135 Leu Arg Phe Lys Ile Leu Arg Ser Leu Gln Ala Phe Val Ala Val Ala

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RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/658,699

DATE: 09/27/2000
 TIME: 09:59:26

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149 <223> OTHER INFORMATION: Description of Unknown Organism: surmised Mus sp.
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152 <221> NAME/KEY: CDS
153 <222> LOCATION: (113)..(700)
155 <220> FEATURE:
156 <221> NAME/KEY: mat_peptide
157 <222> LOCATION: (176)..(700)
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163 Met Leu
164 -20
166 gat tgc aga gca gta ata atg cta tgg ctg ttg ccc tgg gtc act cag 166
167 Asp Cys Arg Ala Val Ile Met Leu Trp Leu Leu Pro Trp Val Thr Gln
168 -15 -10 -5
170 ggc ctg gct gtg cct agg agt agc agt cct gac tgg gct cag tgc cag 214
171 Gly Leu Ala Val Pro Arg Ser Ser Ser Pro Asp Trp Ala Gln Cys Gln
172 -1 1 5 10
174 cag ctc tct cgg aat ctc tgc atg cta gcc tgg aac gca cat gca cca 262
175 Gln Leu Ser Arg Asn Leu Cys Met Leu Ala Trp Asn Ala His Ala Pro
176 15 20 25
178 gcg gga cat atg aat cta cta aga gaa gaa gag gat gaa gag act aaa 310
179 Ala Gly His Met Asn Leu Leu Arg Glu Glu Glu Asp Glu Glu Thr Lys
180 30 35 40 45
182 aat aat gtg ccc cgt atc cag tgt gaa gat ggt tgt gac cca caa gga 358
183 Asn Asn Val Pro Arg Ile Gln Cys Glu Asp Gly Cys Asp Pro Gln Gly
184 50 55 60
186 ctc aag gac aac agc cag ttc tgc ttg caa agg atc cgc caa ggt ctg 406
187 Leu Lys Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile Arg Gln Gly Leu
188 65 70 75
190 gct ttt tat aag cac ctg ctt gac tct gac atc ttc aaa ggg gag cct 454
191 Ala Phe Tyr Lys His Leu Leu Asp Ser Asp Ile Phe Lys Gly Glu Pro
192 80 85 90
194 gct cta ctc cct gat agc ccc atg gag caa ctt cac acc tcc cta cta 502
195 Ala Leu Leu Pro Asp Ser Pro Met Glu Gln Leu His Thr Ser Leu Leu
196 95 100 105
198 gga ctc agc caa ctc ctc cag cca gag gat cac ccc cgg gag acc caa 550
199 Gly Leu Ser Gln Leu Leu Gln Pro Glu Asp His Pro Arg Glu Thr Gln
200 110 115 120 125
202 cag atg ccc agc ctg agt tct agt cag cag tgg cag cgc ccc ctt ctc 598
203 Gln Met Pro Ser Leu Ser Ser Ser Gln Gln Trp Gln Arg Pro Leu Leu

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RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/658,699
 DATE: 09/27/2000
 TIME: 09:59:26

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 Output Set: N:\CRF3\09272000\I658699.raw

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207 Arg Ser Lys Ile Leu Arg Ser Leu Gln Ala Phe Leu Ala Ile Ala Ala
208          145          150          155
210 cgg gtc ttt gcc cac gga gca gca act ctg act gag ccc tta gtg cca 694
211 Arg Val Phe Ala His Gly Ala Ala Thr Leu Thr Glu Pro Leu Val Pro
212          160          165          170
214 aca gct taaggatgcc caggttccca tggctacat gataagacta atctatcagc 750
215 Thr Ala
216          175
218 ccagacatct accagttaat taaccattta ggacttgtgc tgttcttgtt tcgtttgttt 810
220 tgcgtgaagg gcaaggacac cattattaaa gagaaaagaa acaaacccca gagcaggcag 870
222 ctggctagag aaaggagctg gagaagaaga ataaagtctc gagcccttgg ccttggaagc 930
224 gggcaagcag ctgcgtggcc tgaggggaag ggggcggtgg catcgagaaa ctgtgagaaa 990
226 acccagagca tcagaaaaag tgagcccagg ctttggccat tatctgtaag aaaaacaaga 1050
228 aaaggggaac attatacttt cctgggtggc tcagggaaat gtgcagatgc acagtactcc 1110
230 agacagcagc tctgtacctg cctgctctgt ccctcagttc taacagaatc tagtactaa 1170
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240 <220> FEATURE:
241 <223> OTHER INFORMATION: Description of Unknown Organism: surmised Mus sp.
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247 Thr Gln Gly Leu Ala Val Pro Arg Ser Ser Ser Pro Asp Trp Ala Gln
248 -5 -1 1 5 10
250 Cys Gln Gln Leu Ser Arg Asn Leu Cys Met Leu Ala Trp Asn Ala His
251 15 20 25
253 Ala Pro Ala Gly His Met Asn Leu Leu Arg Glu Glu Glu Asp Glu Glu
254 30 35 40
256 Thr Lys Asn Asn Val Pro Arg Ile Gln Cys Glu Asp Gly Cys Asp Pro
257 45 50 55
259 Gln Gly Leu Lys Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile Arg Gln
260 60 65 70 75
262 Gly Leu Ala Phe Tyr Lys His Leu Leu Asp Ser Asp Ile Phe Lys Gly
263 80 85 90
265 Glu Pro Ala Leu Leu Pro Asp Ser Pro Met Glu Gln Leu His Thr Ser
266 95 100 105
268 Leu Leu Gly Leu Ser Gln Leu Leu Gln Pro Glu Asp His Pro Arg Glu
269 110 115 120
271 Thr Gln Gln Met Pro Ser Leu Ser Ser Ser Gln Gln Trp Gln Arg Pro
272 125 130 135
274 Leu Leu Arg Ser Lys Ile Leu Arg Ser Leu Gln Ala Phe Leu Ala Ile
275 140 145 150 155
277 Ala Ala Arg Val Phe Ala His Gly Ala Ala Thr Leu Thr Glu Pro Leu
278 160 165 170

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RAW SEQUENCE LISTING

DATE: 09/27/2000

PATENT APPLICATION: US/09/658,699

TIME: 09:59:26

Input Set : A:\Cpg.pto

Output Set: N:\CRF3\09272000\I658699.raw

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287 <212> TYPE: PRT
288 <213> ORGANISM: Unknown Organism
290 <220> FEATURE:
291 <223> OTHER INFORMATION: Description of Unknown Organism: surmised Sus sp.
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297 Leu Gly Ser Asp Ile Phe Thr Gly Glu Pro Ser Leu His Pro Asp Gly
298      20      25      30
300 Ser Val Gly Gln Leu His Ala Ser Leu Leu Gly Leu Arg Gln Leu Leu
301      35      40      45
303 Gln Pro Glu Gly His His Trp Glu Thr Glu Gln Thr Pro Ser Pro Ser
304      50      55      60
306 Pro Ser Gln Pro Trp Gln Arg Leu Leu Leu Arg Leu Lys Ile Leu Arg
307      65      70      75      80
309 Ser Leu Gln Ala Phe Val Ala Val Ala Ala Arg Val Phe Ala His Gly
310      85      90      95
312 Ala Ala Thr Leu Ser Gln
313      100

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VERIFICATION SUMMARY

DATE: 09/27/2000

PATENT APPLICATION: US/09/658,699

TIME: 09:59:27

Input Set : A:\Cpg.pto

Output Set: N:\CRF3\09272000\I658699.raw

L:15 M:270 C: Current Application Number differs, Replaced Application Number
L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date

RAW SEQUENCE LISTING DATE: 09/22/2000
 PATENT APPLICATION: US/09/658,699 TIME: 15:01:35

Input Set : A:\DX01042X US sequence listing.txt
 Output Set: N:\CRF3\09222000\I658699.raw

Does Not Comply
 Corrected Diskette Needed

pp. 2, 4

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3 <110> APPLICANT: De Waal Malefyt, Rene
4      Kastelein, Robert A.
5      Lira, Sergio A.
6      Narula, Satwant
7      Oppmann, Birgit
8      Rennick, Donna M.
9      Wiekowski, Maria
11 <120> TITLE OF INVENTION: Mammalian Cytokines; Related Reagents and Methods
13 <130> FILE REFERENCE: DX01042X US
C--> 15 <140> CURRENT APPLICATION NUMBER: US/09/658,699
C--> 16 <141> CURRENT FILING DATE: 2000-09-08
18 <150> PRIOR APPLICATION NUMBER: 09/393,090
19 <151> PRIOR FILING DATE: 1999-09-09
21 <150> PRIOR APPLICATION NUMBER: 60/164,616
22 <151> PRIOR FILING DATE: 1999-11-10
24 <160> NUMBER OF SEQ ID NOS: 5
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29 <211> LENGTH: 570
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41 <220> FEATURE:
42 <221> NAME/KEY: mat_peptide
43 <222> LOCATION: (64)..(567)
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50 gct cag ggc aga gct gtg cct ggg ggc agc agc cct gcc tgg act cag 96
51 Ala Gln Gly Arg Ala Val Pro Gly Gly Ser Ser Pro Ala Trp Thr Gln
52 -5 -1 1 5 10
54 tgc cag cag ctt tca cag aag ctc tgc aca ctg gcc tgg agt gca cat 144
55 Cys Gln Gln Leu Ser Gln Lys Leu Cys Thr Leu Ala Trp Ser Ala His
56 15 20 25
58 cca cta gtg gga cac atg gat cta aga gaa gag gga gat gaa gag act 192
59 Pro Leu Val Gly His Met Asp Leu Arg Glu Glu Gly Asp Glu Glu Thr
60 30 35 40
62 aca aat gat gtt ccc cat atc cag tgt gga gat ggc tgt gac ccc caa 240
63 Thr Asn Asp Val Pro His Ile Gln Cys Gly Asp Gly Cys Asp Pro Gln
64 45 50 55
66 gga ctc agg gac aac agt cag ttc tgc ttg caa agg atc cac cag ggt 288

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RAW SEQUENCE LISTING

DATE: 09/22/2000

PATENT APPLICATION: US/09/658,699

TIME: 15:01:35

Input Set : A:\DX01042X US sequence listing.txt

Output Set: N:\CRF3\09222000\I658699.raw

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70 ctg att ttt tat gag aag ctg cta gga tcg gat att ttc aca ggg gag 336
71 Leu Ile Phe Tyr Glu Lys Leu Leu Gly Ser Asp Ile Phe Thr Gly Glu
72 80 85 90
74 cct tct ctg ctc cct gat agc cct gtg gcg cag ctt cat gcc tcc cta 384
75 Pro Ser Leu Leu Pro Asp Ser Pro Val Ala Gln Leu His Ala Ser Leu
76 95 100 105
78 ctg ggc ctc agc caa ctc ctg cag cct gag ggt cac cac tgg gag act 432
79 Leu Gly Leu Ser Gln Leu Leu Gln Pro Glu Gly His His Trp Glu Thr
80 110 115 120
82 cag cag att cca agc ctc agt ccc agc cag cca tgg cag cgt ctc ctt 480
83 Gln Gln Ile Pro Ser Leu Ser Pro Ser Gln Pro Trp Gln Arg Leu Leu
84 125 130 135
86 ctc cgc ttc aaa atc ctt cgc agc ctc cag gcc ttt gtg gct gta gcc 528
87 Leu Arg Phe Lys Ile Leu Arg Ser Leu Gln Ala Phe Val Ala Val Ala
88 140 145 150 155
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92 160 165

95 <210> SEQ ID NO: 2

96 <211> LENGTH: 189

97 <212> TYPE: PRT

98 <213> ORGANISM: Unknown Organism

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99 <223> OTHER INFORMATION: Description of Unknown Organism: surmised Homo

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107 -5 -1 1 5 10

109 Cys Gln Gln Leu Ser Gln Lys Leu Cys Thr Leu Ala Trp Ser Ala His

110 15 20 25

112 Pro Leu Val Gly His Met Asp Leu Arg Glu Glu Gly Asp Glu Glu Thr

113 30 35 40

115 Thr Asn Asp Val Pro His Ile Gln Cys Gly Asp Gly Cys Asp Pro Gln

116 45 50 55

118 Gly Leu Arg Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile His Gln Gly

119 60 65 70 75

121 Leu Ile Phe Tyr Glu Lys Leu Leu Gly Ser Asp Ile Phe Thr Gly Glu

122 80 85 90

124 Pro Ser Leu Leu Pro Asp Ser Pro Val Ala Gln Leu His Ala Ser Leu

125 95 100 105

127 Leu Gly Leu Ser Gln Leu Leu Gln Pro Glu Gly His His Trp Glu Thr

128 110 115 120

130 Gln Gln Ile Pro Ser Leu Ser Pro Ser Gln Pro Trp Gln Arg Leu Leu

131 125 130 135

133 Leu Arg Phe Lys Ile Leu Arg Ser Leu Gln Ala Phe Val Ala Val Ala

134 140 145 150 155

<220> feature is
required here.

RAW SEQUENCE LISTING DATE: 09/22/2000
 PATENT APPLICATION: US/09/658,699 TIME: 15:01:35

Input Set : A:\DX01042X US sequence listing.txt
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150 <221> NAME/KEY: CDS
151 <222> LOCATION: (113)..(700)
153 <220> FEATURE:
154 <221> NAME/KEY: mat_peptide
155 <222> LOCATION: (176)..(700)
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161                               Met Leu
162                               -20
164 gat tgc aga gca gta ata atg cta tgg ctg ttg ccc tgg gtc act cag 166
165 Asp Cys Arg Ala Val Ile Met Leu Trp Leu Leu Pro Trp Val Thr Gln
166                               -15           -10           -5
168 ggc ctg gct gtg cct agg agt agc agt cct gac tgg gct cag tgc cag 214
169 Gly Leu Ala Val Pro Arg Ser Ser Ser Pro Asp Trp Ala Gln Cys Gln
170                               -1    1           5           10
172 cag etc tct cgg aat etc tgc atg cta gcc tgg aac gca cat gca cca 262
173 Gln Leu Ser Arg Asn Leu Cys Met Leu Ala Trp Asn Ala His Ala Pro
174                               15           20           25
176 gcg gga cat atg aat cta cta aga gaa gaa gag gat gaa gag act aaa 310
177 Ala Gly His Met Asn Leu Leu Arg Glu Glu Glu Asp Glu Glu Thr Lys
178 30           35           40           45
180 aat aat gtg ccc cgt atc cag tgt gaa gat ggt tgt gac cca caa gga 358
181 Asn Asn Val Pro Arg Ile Gln Cys Glu Asp Gly Cys Asp Pro Gln Gly
182                               50           55           60
184 etc aag gac aac agc cag ttc tgc ttg caa agg atc cgc caa ggt ctg 406
185 Leu Lys Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile Arg Gln Gly Leu
186                               65           70           75
188 gct ttt tat aag cac ctg ctt gac tct gac atc ttc aaa ggg gag cct 454
189 Ala Phe Tyr Lys His Leu Leu Asp Ser Asp Ile Phe Lys Gly Glu Pro
190                               80           85           90
192 gct cta etc cct gat agc ccc atg gag caa ctt cac acc tcc cta cta 502
193 Ala Leu Leu Pro Asp Ser Pro Met Glu Gln Leu His Thr Ser Leu Leu
194                               95           100          105
196 gga etc agc caa etc ctc cag cca gag gat cac ccc cgg gag acc caa 550
197 Gly Leu Ser Gln Leu Leu Gln Pro Glu Asp His Pro Arg Glu Thr Gln
198 110          115          120          125
200 cag atg ccc agc ctg agt tct agt cag cag tgg cag cgc ccc ctt etc 598
201 Gln Met Pro Ser Leu Ser Ser Ser Gln Gln Trp Gln Arg Pro Leu Leu
202                               130          135          140

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RAW SEQUENCE LISTING

DATE: 09/22/2000

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TIME: 15:01:35

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Output Set: N:\CRF3\09222000\I658699.raw

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206 145 150 155
208 cgg gtc ttt gcc cac gga gca gca act ctg act gag ccc tta gtg cca 694
209 Arg Val Phe Ala His Gly Ala Ala Thr Leu Thr Glu Pro Leu Val Pro
210 160 165 170
212 aca gct taaggatgcc caggttccca tggctaccat gataagacta atctatcagc 750
213 Thr Ala
214 175
216 ccagacatct accagttaat taaccatta ggacttctgc tgttcttgtt tcgtttgttt 810
218 tgcgtgaagg gcaaggacac cattattaaa gagaaaagaa acaaacccca gagcaggcag 870
220 ctggctagag aaaggagctg gagaagaaga ataaagtctc gagcccttgg ccttggaagc 930
222 gggcaagcag ctgcgtggcc tgagggaag ggggcggtgg catcgagaaa ctgtgagaaa 990
224 acccagagca tcagaaaaag tgagcccagg ctttggccat tatctgtaag aaaaacaaga 1050
226 aaaggggaac attatacttt cctgggtggc tcagggaaat gtgcagatgc acagtactcc 1110
228 agacagcagc tctgtacctg cctgctctgt ccttcagttc taacagaatc tagtcactaa 1170
230 gaactaacag gactaccaat acgaactgac aaa 1203
233 <210> SEQ ID NO: 4
234 <211> LENGTH: 196
235 <212> TYPE: PRT
236 <213> ORGANISM: Unknown Organism
W--> 237 <220> FEATURE:
237 <223> OTHER INFORMATION: Description of Unknown Organism: surmised Mus sp.
239 <400> SEQUENCE: 4
240 Met Leu Asp Cys Arg Ala Val Ile Met Leu Trp Leu Leu Pro Trp Val
241 -20 -15 -10
243 Thr Gln Gly Leu Ala Val Pro Arg Ser Ser Ser Pro Asp Trp Ala Gln
244 -5 -1 1 5 10
246 Cys Gln Gln Leu Ser Arg Asn Leu Cys Met Leu Ala Trp Asn Ala His
247 15 20 25
249 Ala Pro Ala Gly His Met Asn Leu Leu Arg Glu Glu Glu Asp Glu Glu
250 30 35 40
252 Thr Lys Asn Asn Val Pro Arg Ile Gln Cys Glu Asp Gly Cys Asp Pro
253 45 50 55
255 Gln Gly Leu Lys Asp Asn Ser Gln Phe Cys Leu Gln Arg Ile Arg Gln
256 60 65 70 75
258 Gly Leu Ala Phe Tyr Lys His Leu Leu Asp Ser Asp Ile Phe Lys Gly
259 80 85 90
261 Glu Pro Ala Leu Leu Pro Asp Ser Pro Met Glu Gln Leu His Thr Ser
262 95 100 105
264 Leu Leu Gly Leu Ser Gln Leu Leu Gln Pro Glu Asp His Pro Arg Glu
265 110 115 120
267 Thr Gln Gln Met Pro Ser Leu Ser Ser Ser Gln Gln Trp Gln Arg Pro
268 125 130 135
270 Leu Leu Arg Ser Lys Ile Leu Arg Ser Leu Gln Ala Phe Leu Ala Ile
271 140 145 150 155
273 Ala Ala Arg Val Phe Ala His Gly Ala Ala Thr Leu Thr Glu Pro Leu
274 160 165 170
276 Val Pro Thr Ala

<220> feature
is required here

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/658,699
DATE: 09/22/2000
TIME: 15:01:35

Input Set : A:\DX01042X US sequence listing.txt
Output Set: N:\CRF3\09222000\I658699.raw

277 175
281 <210> SEQ ID NO: 5
282 <211> LENGTH: 102
283 <212> TYPE: PRT
284 <213> ORGANISM: Unknown Organism
286 <220> FEATURE:
287 <223> OTHER INFORMATION: Description of Unknown Organism: surmised Sus sp.
289 <400> SEQUENCE: 5
290 Ser Cys Leu Gln Arg Ile His Gln Gly Leu Val Phe Tyr Glu Lys Leu
291 1 5 10 15
293 Leu Gly Ser Asp Ile Phe Thr Gly Glu Pro Ser Leu His Pro Asp Gly
294 20 25 30
296 Ser Val Gly Gln Leu His Ala Ser Leu Leu Gly Leu Arg Gln Leu Leu
297 35 40 45
299 Gln Pro Glu Gly His His Trp Glu Thr Glu Gln Thr Pro Ser Pro Ser
300 50 55 60
302 Pro Ser Gln Pro Trp Gln Arg Leu Leu Leu Arg Leu Lys Ile Leu Arg
303 65 70 75 80
305 Ser Leu Gln Ala Phe Val Ala Val Ala Ala Arg Val Phe Ala His Gly
306 85 90 95
308 Ala Ala Thr Leu Ser Gln
309 100

VERIFICATION SUMMARY

DATE: 09/22/2000

PATENT APPLICATION: US/09/658,699

TIME: 15:01:36

Input Set : A:\DX01042X US sequence listing.txt

Output Set: N:\CRF3\09222000\I658699.raw

L:15 M:270 C: Current Application Number differs, Replaced Application Number

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date

L:99 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:237 M:258 W: Mandatory Feature missing, <220> FEATURE: